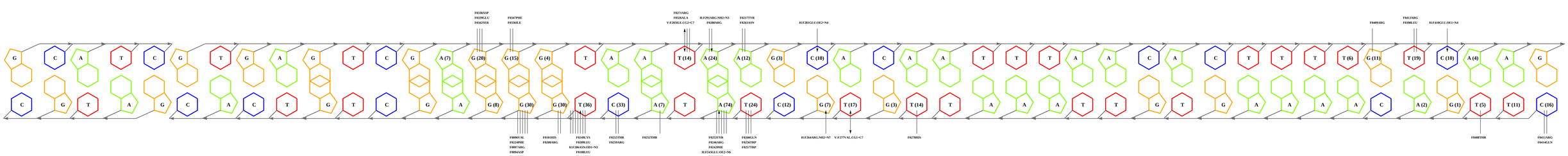
F013BASP
F013SGLU
F034ZSER
F034?PHE
F035ILE
F027SARG
F028ALA
V:F28ILE,CG2>C?
H:F29ARG,NH2>N3
F028BARG
F021?TYR
F026ASN
H:F28GLU,OE2>N4
F009SARG
F010HLEU
F041SARG
F039HLEU
H:F41GLU,OE1>N4
G C A T C G T G A G T C G A (7) G (20) G (15) G (4) T A A T (14) A (24) A (12) G (3) C (10) A C A A T T T A A C A C T T T T (6) G (11) T (19) C (10) A (4) A G
C G T A G C A C T G T C G A G (8) G (30) G (30) T (36) C (33) A (7) T A (74) T (24) C (12) G (7) T (17) G (3) T (14) T A A A T T G T G A A A A C A (2) G (1) T (5) T (11) C (16)
F009SVAL
F022ARSE
F009TARG
F009AASP
F010HHIS
F028BARG
F024PLYS
F020PLEU
H:F29GASN,OD1>N3
F010HLEU
F025TTHR
F025DARG
F025?THR
F022TTYR
F020AARG
F024ZPHE
H:F24GGLU,OE2>N4
F026SGLN
F026TRP
F025TTRP
H:F26ARG,NH2>N7
V:F27?VAL,CG1>C?
F027HHIS
F040HTHR
F041ARSG
F041?GLN